

Figure 1-A

>2720879 GGCCACCGGG ACTTCAGTGT CTCCTCCATC CCAGGAGCGC AGTGGCCACT
>1362407 ACCGGG ACTTCAGTGT CTCCTCCATC CCAGGAGCGC AGTGGCCACT
>1362407IH ACCGGG ACTTCAGTGT CTCCTCCATC CCAGGAGCGC AGTGGCCACT
>1512552 CAGGAGCGC AGTGGCCACT
>1512552IH CAGGAGCGC AGTGGCCACT
Consensus GGCCACCGGG ACTTCAGTGT CTCCTCCATC CCAGGAGCGC AGTGGCCACT

>2720879 ATGGGGTCTG GGCTGCCCT TGTCCTCCTC TTGACCCCTCC TTGGCAGCTC
>1362407 ATGGGGTCTG GGCTGCCCT TGTCCTCCTC TTGACCCCTCC TTGGCAGCTC
>1362407IH ATGGGGTCTG GGCTGCCCT TGTCCTCCTC TTGACCCCTCC TTGGCAGCTC
>1512552 ATGGGGTCTG GGCTGCCCT TGTCCTCCTC TTGACCCCTCC TTGGCAGCTC
>1512552IH ATGGGGTCTG GGCTGCCCT TGTCCTCCTC TTGACCCCTCC TTGGCAGCTC
Consensus ATGGGGTCTG GGCTGCCCT TGTCCTCCTC TTGACCCCTCC TTGGCAGCTC

>2720879 ACATGGAACA GGGCCGGGT TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
>1362407 ACATGGAACA GGGCCGGGT TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
>1362407IH ACATGGAACA GGGCCGGGT TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
>1512552 ACATGGAACA GGGCCGGGT TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
>1512552IH ACATGGAACA GGGCCGGGT TGACTTTGCA ACTGAAGCTG AAGGAGTCTT
Consensus ACATGGAACA GGGCCGGGT TGACTTTGCA ACTGAAGCTG AAGGAGTCTT

>2720879 TTCTGACAAA TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAN
>1362407 TTCTGACAAA TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG
>1362407IH TTCTGACAAA TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG
>1512552 TTCTGACAAG TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG
>1512552IH TTCTGACAAG TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG
Consensus TTCTGACAAA TTCCTCCTAT GAGTCCAGCT TCCTGGAATT GCTTGAAAAG

>2720879 :TCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA
>1362407 :TCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCA
>1362407IH CTCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA CCCTCCACCA
>1512552 CTCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA CCCTCCACCA
>1512552IH CTCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA CCCTCCACCA
Consensus CTCTGCCTCC TCCTCCATCT CCCTTCAGGG ACCAGCGTCA CCCTCCACCA

>1362407IH TGCAAGATCT CAACACCATG TTGTCTGCAA CACATGACAG CCATTGAAGC
>1512552IH TGCAAGATCT CAACACCATG TTGTCTGCAA CACATGACAG CCATTGAAGC
>g727537 T CAACACCATG TTGTCTGCAA CACATGACAG CCATTGAAGC
Consensus TGCAAGATCT CAACACCATG TTGTCTGCAA CACATGACAG CCATTGAAGC

>1362407IH CTGTGTCCTT CTTGGCCCCGG GCTTTTGGGC CGGGGATGCA GGAGGCAGGC
>1512552IH CTGTGTCCTT CTTGGCCCCGG GCTTTTGGGC CGGGGATGCA GGAGGCAGGC
>g727537 CTGTGTCCTT CTTGGCCCCGG GCTTTTNGGC GGGGAATGCA GGAGGCAGGC
Consensus CTGTGTCCTT CTTGGCCCCGG GCTTTTGGGC CGGGGATGCA GGAGGCAGGC

>1362407IH CCCGACCCCTG TCTTTCAGCA GGCCCCCACC CTCCTGAGTG GCAATAAATA
>1512552IH CCCGACCCCTG TCTTTCAGCA GGCCCCCACC CTCCTGAGTG GCAATAAATA
>g727537 CCCGACCCCTN TCTTTCAGCA GGCCCCCACC CTNCTGAGTN GCAATAAATA
Consensus CCCGACCCCTG TCTTTCAGCA GGCCCCCACC CTCCTGAGTG GCAATAAATA

Figure 1-B

>1362407IH	AAATTCGGTA TGCTTGA
>1512552IH	AAATTCGGTA TGCTGAATT
>g727537	AAATTCGGTA TGCTGAATT C AATA
Consensus	AAATTCGGTA TGCTGAATT C AATA

Figure 2

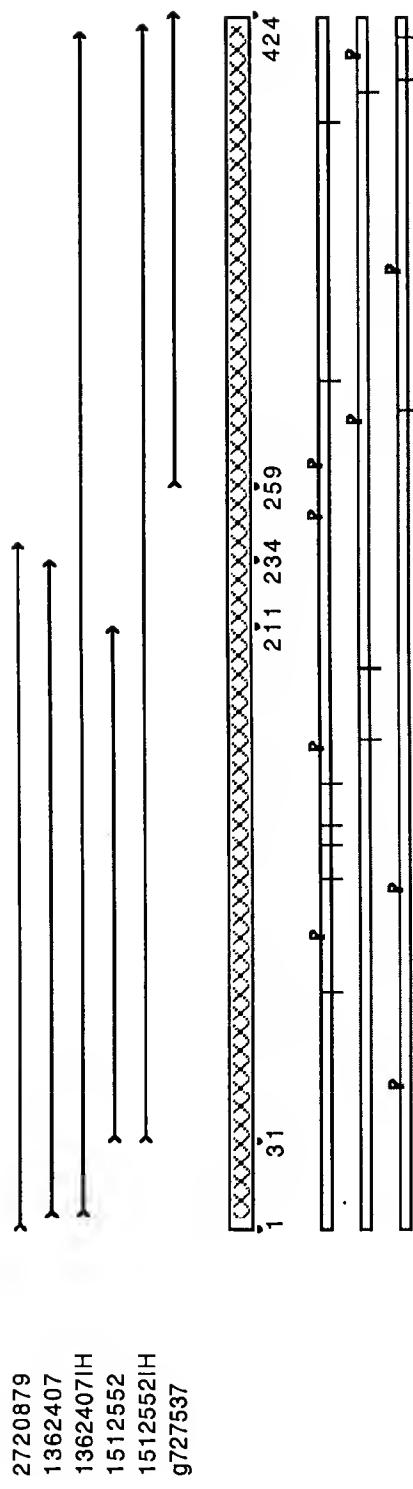
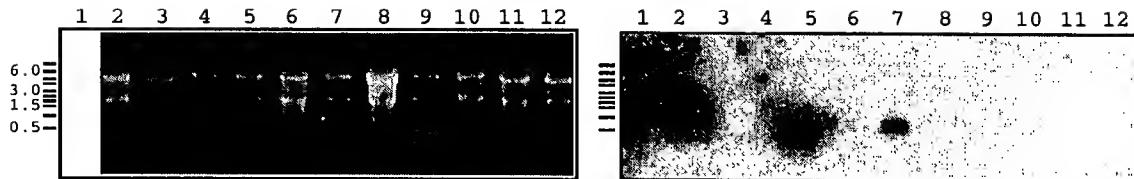


Figure 3



Lane	Tissue	Lane	Tissue
1	Bladder	7	Lung
2	Brain	8	Ovary
3	Breast	9	Placenta
4	Colon	10	Prostate
5	Kidney	11	Spleen
6	Liver	12	Testis